

EAST Search History

Ref #	Hits	Search Query	DBs	Default Operator	Plurals	Time Stamp
L5	270400	transcription or translation or protein synthesis	US-PGPUB; USPAT	ADJ	OFF	2006/10/06 16:30
L6	9992	"1" near5 (in vitro or cell free) or itt	US-PGPUB; USPAT	ADJ	OFF	2006/10/06 16:31
L7	369	recBC\$ or exov or exo adj v	US-PGPUB; USPAT	ADJ	OFF	2006/10/06 16:31
(L8)	26	6 and 7	US-PGPUB; USPAT	ADJ	OFF	2006/10/06 16:31

* * * * * STN Columbus * * * * *

FILE 'HOME' ENTERED AT 16:40:05 ON 06 OCT 2006

=> fil .bec

COST IN U.S. DOLLARS

SINCE FILE	TOTAL
ENTRY	SESSION
0.21	0.21

FULL ESTIMATED COST

FILES 'MEDLINE, SCISEARCH, LIFESCI, BIOTECHDS, BIOSIS, EMBASE, HCAPLUS, NTIS, ESBIODBASE, BIOTECHNO, WPIDS' ENTERED AT 16:40:18 ON 06 OCT 2006
ALL COPYRIGHTS AND RESTRICTIONS APPLY. SEE HELP USAGETERMS FOR DETAILS.

11 FILES IN THE FILE LIST

=> s transcription or translation or protein synthesis

FILE 'MEDLINE'

281709 TRANSCRIPTION
45558 TRANSLATION
1575820 PROTEIN
433491 SYNTHESIS
51283 PROTEIN SYNTHESIS
(PROTEIN(W) SYNTHESIS)

L1 354106 TRANSCRIPTION OR TRANSLATION OR PROTEIN SYNTHESIS

FILE 'SCISEARCH'

223168 TRANSCRIPTION
54572 TRANSLATION
1331797 PROTEIN
772277 SYNTHESIS
44136 PROTEIN SYNTHESIS
(PROTEIN(W) SYNTHESIS)

L2 302012 TRANSCRIPTION OR TRANSLATION OR PROTEIN SYNTHESIS

FILE 'LIFESCI'

115478 TRANSCRIPTION
25086 TRANSLATION
514071 "PROTEIN"
111352 "SYNTHESIS"
16572 PROTEIN SYNTHESIS
("PROTEIN" (W) "SYNTHESIS")

L3 145350 TRANSCRIPTION OR TRANSLATION OR PROTEIN SYNTHESIS

FILE 'BIOTECHDS'

20252 TRANSCRIPTION
6231 TRANSLATION
152563 PROTEIN
24620 SYNTHESIS
1436 PROTEIN SYNTHESIS
(PROTEIN(W) SYNTHESIS)

L4 25096 TRANSCRIPTION OR TRANSLATION OR PROTEIN SYNTHESIS

FILE 'BIOSIS'

242977 TRANSCRIPTION
54523 TRANSLATION
1600576 PROTEIN
566268 SYNTHESIS
73491 PROTEIN SYNTHESIS
(PROTEIN(W) SYNTHESIS)

L5 344615 TRANSCRIPTION OR TRANSLATION OR PROTEIN SYNTHESIS

FILE 'EMBASE'

271541 TRANSCRIPTION
45716 TRANSLATION

1547249 "PROTEIN"
 571999 "SYNTHESIS"
 85054 PROTEIN SYNTHESIS
 ("PROTEIN" (W) "SYNTHESIS")
 L6 371226 TRANSCRIPTION OR TRANSLATION OR PROTEIN SYNTHESIS

FILE 'HCAPLUS'
 327803 TRANSCRIPTION
 495728 TRANSLATION
 1921265 PROTEIN
 1271581 SYNTHESIS
 72621 PROTEIN SYNTHESIS
 (PROTEIN (W) SYNTHESIS)
 L7 856750 TRANSCRIPTION OR TRANSLATION OR PROTEIN SYNTHESIS

FILE 'NTIS'
 2045 TRANSCRIPTION
 64328 TRANSLATION
 13503 PROTEIN
 35555 SYNTHESIS
 615 PROTEIN SYNTHESIS
 (PROTEIN (W) SYNTHESIS)
 L8 66785 TRANSCRIPTION OR TRANSLATION OR PROTEIN SYNTHESIS

FILE 'ESBIOBASE'
 130298 TRANSCRIPTION
 23657 TRANSLATION
 669718 PROTEIN
 171104 SYNTHESIS
 41400 PROTEIN SYNTHESIS
 (PROTEIN (W) SYNTHESIS)
 L9 173078 TRANSCRIPTION OR TRANSLATION OR PROTEIN SYNTHESIS

FILE 'BIOTECHNO'
 160885 TRANSCRIPTION
 25603 TRANSLATION
 623255 PROTEIN
 144368 SYNTHESIS
 32236 PROTEIN SYNTHESIS
 (PROTEIN (W) SYNTHESIS)
 L10 200107 TRANSCRIPTION OR TRANSLATION OR PROTEIN SYNTHESIS

FILE 'WPIDS'
 15483 TRANSCRIPTION
 29849 TRANSLATION
 144848 PROTEIN
 90084 SYNTHESIS
 1285 PROTEIN SYNTHESIS
 (PROTEIN (W) SYNTHESIS)
 L11 44338 TRANSCRIPTION OR TRANSLATION OR PROTEIN SYNTHESIS

TOTAL FOR ALL FILES
 L12 2883463 TRANSCRIPTION OR TRANSLATION OR PROTEIN SYNTHESIS

=> s l12(5a) (in vitro or cell free) or itt
 FILE 'MEDLINE'
 11123291 IN
 845983 VITRO
 845708 IN VITRO
 (IN (W) VITRO)
 2055382 CELL
 487418 FREE
 32058 CELL FREE
 (CELL (W) FREE)
 16244 L1 (5A) (IN VITRO OR CELL FREE)

1030 ITT
 L13 17271 L1 (5A) (IN VITRO OR CELL FREE) OR ITT

 FILE 'SCISEARCH'
 14989233 IN
 484855 VITRO
 481422 IN VITRO
 (IN(W)VITRO)
 1595227 CELL
 627623 FREE
 17350 CELL FREE
 (CELL(W)FREE)
 9925 L2 (5A) (IN VITRO OR CELL FREE)
 1166 ITT
 L14 11088 L2 (5A) (IN VITRO OR CELL FREE) OR ITT

 FILE 'LIFESCI'
 202565 IN VITRO
 ("VITRO")
 546120 "CELL"
 103299 "FREE"
 10243 CELL FREE
 ("CELL" (W) "FREE")
 9316 L3 (5A) (IN VITRO OR CELL FREE)
 112 ITT
 L15 9427 L3 (5A) (IN VITRO OR CELL FREE) OR ITT

 FILE 'BIOTECHDS'
 29429 IN VITRO
 (VITRO)
 170601 CELL
 28129 FREE
 4648 CELL FREE
 (CELL(W)FREE)
 1584 L4 (5A) (IN VITRO OR CELL FREE)
 6 ITT
 L16 1590 L4 (5A) (IN VITRO OR CELL FREE) OR ITT

 FILE 'BIOSIS'
 681157 IN VITRO
 (VITRO)
 3791623 CELL
 484410 FREE
 27084 CELL FREE
 (CELL(W)FREE)
 18920 L5 (5A) (IN VITRO OR CELL FREE)
 923 ITT
 L17 19840 L5 (5A) (IN VITRO OR CELL FREE) OR ITT

 FILE 'EMBASE'
 8995168 "IN"
 985223 "VITRO"
 984960 IN VITRO
 ("IN" (W) "VITRO")
 2719416 "CELL"
 406882 "FREE"
 19029 CELL FREE
 ("CELL" (W) "FREE")
 13070 L6 (5A) (IN VITRO OR CELL FREE)
 1059 ITT
 L18 14126 L6 (5A) (IN VITRO OR CELL FREE) OR ITT

 FILE 'HCAPLUS'
 641187 IN VITRO
 (VITRO)

2118574 CELL
 1269684 FREE
 36275 CELL FREE
 (CELL(W) FREE)
 21844 L7 (5A) (IN VITRO OR CELL FREE)
 747 ITT
 L19 22589 L7 (5A) (IN VITRO OR CELL FREE) OR ITT

FILE 'NTIS'

 1785086 IN
 9133 VITRO
 9044 IN VITRO
 (IN(W) VITRO)
 50856 CELL
 61685 FREE
 354 CELL FREE
 (CELL(W) FREE)
 132 L8 (5A) (IN VITRO OR CELL FREE)
 166 ITT
 L20 298 L8 (5A) (IN VITRO OR CELL FREE) OR ITT

FILE 'ESBIOBASE'

 2979532 IN
 220542 VITRO
 220312 IN VITRO
 (IN(W) VITRO)
 803565 CELL
 157557 FREE
 7481 CELL FREE
 (CELL(W) FREE)
 6190 L9 (5A) (IN VITRO OR CELL FREE)
 446 ITT
 L21 6633 L9 (5A) (IN VITRO OR CELL FREE) OR ITT

FILE 'BIOTECHNO'

 1588351 IN
 253158 VITRO
 253028 IN VITRO
 (IN(W) VITRO)
 822843 CELL
 81349 FREE
 9281 CELL FREE
 (CELL(W) FREE)
 9667 L10 (5A) (IN VITRO OR CELL FREE)
 93 ITT
 L22 9758 L10 (5A) (IN VITRO OR CELL FREE) OR ITT

FILE 'WPIDS'

 26891 IN VITRO
 (VITRO)
 419630 CELL
 532886 FREE
 2720 CELL FREE
 (CELL(W) FREE)
 814 L11 (5A) (IN VITRO OR CELL FREE)
 38 ITT
 L23 852 L11 (5A) (IN VITRO OR CELL FREE) OR ITT

TOTAL FOR ALL FILES

L24 113472 L12 (5A) (IN VITRO OR CELL FREE) OR ITT

=> s (nuclease# or ribonuclease# or deoxyribonuclease# or rnase## or
 dnase##) (4a) inhibit?

FILE 'MEDLINE'

 16567 NUCLEASE#

28595 RIBONUCLEASE#
25717 DEOXYRIBONUCLEASE#
14251 RNASE##
11323 DNASE##
1286323 INHIBIT?
L25 2217 (NUCLEASE# OR RIBONUCLEASE# OR DEOXYRIBONUCLEASE# OR RNASE## OR
DNASE##) (4A) INHIBIT?

FILE 'SCISEARCH'

11723 NUCLEASE#
12866 RIBONUCLEASE#
1938 DEOXYRIBONUCLEASE#
12477 RNASE##
8762 DNASE##
1077167 INHIBIT?
L26 1645 (NUCLEASE# OR RIBONUCLEASE# OR DEOXYRIBONUCLEASE# OR RNASE## OR
DNASE##) (4A) INHIBIT?

FILE 'LIFESCI'

7855 NUCLEASE#
5887 RIBONUCLEASE#
4882 DEOXYRIBONUCLEASE#
7390 RNASE##
6070 DNASE##
346647 INHIBIT?
L27 938 (NUCLEASE# OR RIBONUCLEASE# OR DEOXYRIBONUCLEASE# OR RNASE## OR
DNASE##) (4A) INHIBIT?

FILE 'BIOTECHDS'

2378 NUCLEASE#
614 RIBONUCLEASE#
125 DEOXYRIBONUCLEASE#
1127 RNASE##
634 DNASE##
60638 INHIBIT?
L28 326 (NUCLEASE# OR RIBONUCLEASE# OR DEOXYRIBONUCLEASE# OR RNASE## OR
DNASE##) (4A) INHIBIT?

FILE 'BIOSIS'

24658 NUCLEASE#
8522 RIBONUCLEASE#
1292 DEOXYRIBONUCLEASE#
27417 RNASE##
17234 DNASE##
1377832 INHIBIT?
L29 2822 (NUCLEASE# OR RIBONUCLEASE# OR DEOXYRIBONUCLEASE# OR RNASE## OR
DNASE##) (4A) INHIBIT?

FILE 'EMBASE'

11623 NUCLEASE#
15685 RIBONUCLEASE#
8581 DEOXYRIBONUCLEASE#
12358 RNASE##
9665 DNASE##
1177306 INHIBIT?
L30 1716 (NUCLEASE# OR RIBONUCLEASE# OR DEOXYRIBONUCLEASE# OR RNASE## OR
DNASE##) (4A) INHIBIT?

FILE 'HCAPLUS'

25579 NUCLEASE#
13676 RIBONUCLEASE#
38290 RNASE
43291 RIBONUCLEASE#
(RIBONUCLEASE# OR RNASE)
3737 DEOXYRIBONUCLEASE#

19911 DNASE
 21608 DEOXYRIBONUCLEASE#
 (DEOXYRIBONUCLEASE# OR DNASE)
 39138 RNASE##
 20513 DNASE##
 1867961 INHIBIT?
 L31 5276 (NUCLEASE# OR RIBONUCLEASE# OR DEOXYRIBONUCLEASE# OR RNASE## OR
 DNASE##) (4A) INHIBIT?

FILE 'NTIS'

206 NUCLEASE#
 193 RIBONUCLEASE#
 47 DEOXYRIBONUCLEASE#
 99 RNASE##
 65 DNASE##
 21289 INHIBIT?
 L32 27 (NUCLEASE# OR RIBONUCLEASE# OR DEOXYRIBONUCLEASE# OR RNASE## OR
 DNASE##) (4A) INHIBIT?

FILE 'ESBIOBASE'

4807 NUCLEASE#
 5821 RIBONUCLEASE#
 524 DEOXYRIBONUCLEASE#
 7487 RNASE##
 4889 DNASE##
 483136 INHIBIT?
 L33 873 (NUCLEASE# OR RIBONUCLEASE# OR DEOXYRIBONUCLEASE# OR RNASE## OR
 DNASE##) (4A) INHIBIT?

FILE 'BIOTECHNO'

7602 NUCLEASE#
 7816 RIBONUCLEASE#
 4089 DEOXYRIBONUCLEASE#
 8055 RNASE##
 6433 DNASE##
 301415 INHIBIT?
 L34 886 (NUCLEASE# OR RIBONUCLEASE# OR DEOXYRIBONUCLEASE# OR RNASE## OR
 DNASE##) (4A) INHIBIT?

FILE 'WPIDS'

2062 NUCLEASE#
 901 RIBONUCLEASE#
 1200 RNASE
 1947 RIBONUCLEASE#
 (RIBONUCLEASE# OR RNASE)
 244 DEOXYRIBONUCLEASE#
 548 DNASE
 755 DEOXYRIBONUCLEASE#
 (DEOXYRIBONUCLEASE# OR DNASE)
 1348 RNASE##
 622 DNASE##
 257362 INHIBIT?
 L35 395 (NUCLEASE# OR RIBONUCLEASE# OR DEOXYRIBONUCLEASE# OR RNASE## OR
 DNASE##) (4A) INHIBIT?

TOTAL FOR ALL FILES

L36 17121 (NUCLEASE# OR RIBONUCLEASE# OR DEOXYRIBONUCLEASE# OR RNASE## OR
 DNASE##) (4A) INHIBIT?

=> s 124 and 136

FILE 'MEDLINE'

L37 85 L13 AND L25

FILE 'SCISEARCH'

L38 47 L14 AND L26

FILE 'LIFESCI'
L39 46 L15 AND L27

FILE 'BIOTECHDS'
L40 26 L16 AND L28

FILE 'BIOSIS'
L41 107 L17 AND L29

FILE 'EMBASE'
L42 63 L18 AND L30

FILE 'HCAPLUS'
L43 178 L19 AND L31

FILE 'NTIS'
L44 0 L20 AND L32

FILE 'ESBIOBASE'
L45 33 L21 AND L33

FILE 'BIOTECHNO'
L46 46 L22 AND L34

FILE 'WPIDS'
L47 18 L23 AND L35

TOTAL FOR ALL FILES
L48 649 L24 AND L36

=> s l48 not 2002-2006/py

FILE 'MEDLINE'
2844973 2002-2006/PY
(20020000-20069999/PY)
L49 75 L37 NOT 2002-2006/PY

FILE 'SCISEARCH'
5243521 2002-2006/PY
(20020000-20069999/PY)
L50 36 L38 NOT 2002-2006/PY

FILE 'LIFESCI'
500908 2002-2006/PY
L51 39 L39 NOT 2002-2006/PY

FILE 'BIOTECHDS'
122693 2002-2006/PY
L52 1 L40 NOT 2002-2006/PY

FILE 'BIOSIS'
2511100 2002-2006/PY
L53 98 L41 NOT 2002-2006/PY

FILE 'EMBASE'
2473613 2002-2006/PY
L54 55 L42 NOT 2002-2006/PY

FILE 'HCAPLUS'
5401054 2002-2006/PY
L55 152 L43 NOT 2002-2006/PY

FILE 'NTIS'
71750 2002-2006/PY
L56 0 L44 NOT 2002-2006/PY

FILE 'ESBIOBASE'
1474871 2002-2006/PY
L57 25 L45 NOT 2002-2006/PY

FILE 'BIOTECHNO'
244553 2002-2006/PY
L58 42 L46 NOT 2002-2006/PY

FILE 'WPIDS'
4666175 2002-2006/PY
L59 8 L47 NOT 2002-2006/PY

TOTAL FOR ALL FILES
L60 531 L48 NOT 2002-2006/PY

=> dup rem l60
PROCESSING COMPLETED FOR L60
L61 192 DUP REM L60 (339 DUPLICATES REMOVED)

=> d tot

L61 ANSWER 1 OF 192 WPIDS COPYRIGHT 2006 THE THOMSON CORP on STN
TI New nucleic acid encoding mammalian capping enzyme, useful for catalyzing
formation of RNA 5'-terminal GpppN cap complex and in complementation
assay to identify and/or monitor genetic defect in capping pathway.
PI US 6312926 B1 20011106 (200203)* 47 C12P019-34
IN MALDONADO, E; PILLUTLA, R; REINBERG, D; SHATKIN, A J; YUE, Z

L61 ANSWER 2 OF 192 WPIDS COPYRIGHT 2006 THE THOMSON CORP on STN
TI Generating a complete full-length cDNA library from single cells for use
in gene chip technology, involves reverse transcribing intracellular
mRNAs, adding polynucleotide tail and amplifying formed cDNAs.
PI US 6197554 B1 20010306 (200125)* 11 C12P019-34
IN CHUONG, C; LIN, S; YING, S

L61 ANSWER 3 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN
TI Executioner caspase-3, -6, and -7 perform distinct, non-redundant roles
during the demolition phase of apoptosis
SO Journal of Biological Chemistry (2001), 276(10), 7320-7326
CODEN: JBCHA3; ISSN: 0021-9258
AU Slee, Elizabeth A.; Adrain, Colin; Martin, Seamus J.
AN 2001:276527 HCAPLUS
DN 134:349517

L61 ANSWER 4 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN
TI Quantification of in vitro retroviral replication using a one-tube
real-time RT-PCR system incorporating direct RNA preparation
SO Journal of Virological Methods (2001), 91(2), 149-155
CODEN: JVMEDH; ISSN: 0166-0934
AU Bisset, L. R.; Bosbach, S.; Tomasik, Z.; Lutz, H.; Schupbach, J.; Boni, J.
AN 2001:74729 HCAPLUS
DN 135:221944

L61 ANSWER 5 OF 192 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on
STN DUPLICATE 1
TI Isolation of a novel deoxyribonuclease with antifungal activity from
Asparagus officinalis seeds.
SO Biochemical and Biophysical Research Communications, (November 23, 2001)
Vol. 289, No. 1, pp. 120-124. print.
CODEN: BBRCA9. ISSN: 0006-291X.
AU Wang, Hexiang; Ng, T. B. [Reprint author]
AN 2002:40724 BIOSIS

L61 ANSWER 6 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 2

TI RNA polymerase chain reaction for generating amplified mRNAs from limited mRNAs
 SO PCT Int. Appl., 31 pp.
 CODEN: PIXXD2
 IN Lin, Shi-Lung; Ying, Shao-Yao; Chuong, Cheng-Ming; Widelitz, Randall Bruce
 AN 2000:881343 HCAPLUS
 DN 134:37905

	PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
PI	WO 2000075356	A1	20001214	WO 1999-US12461	19990604
	W: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, CZ, DE, DE, DK, DK, EE, EE, ES, FI, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SK, SL, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM				
	RW: GH, GM, KE, LS, MW, SD, SL, SZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG				
	AU 9943323	A1	20001228	AU 1999-43323	19990604

L61 ANSWER 7 OF 192 MEDLINE on STN DUPLICATE 3
 TI Specific chaperone-like activity of inhibitor of caspase-activated DNase for caspase-activated DNase.
 SO The Journal of biological chemistry, (2000 Mar 17) Vol. 275, No. 11, pp. 8091-6.
 Journal code: 2985121R. ISSN: 0021-9258.
 AU Sakahira H; Iwamatsu A; Nagata S
 AN 2000179917 MEDLINE

L61 ANSWER 8 OF 192 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN
 AN 2000214279 ESBIIOBASE
 TI First demonstration of lactoribonuclease, a ribonuclease from bovine milk with similarity to bovine pancreatic ribonuclease
 AU Ye X.Y.; Ng T.B.
 CS X.Y. Ye, Department of Biochemistry, Faculty of Medicine, Chinese University of Hong Kong, Shatin, New Territories, Hong Kong.
 SO Life Sciences, (08 SEP 2000), 67/16 (2025-2032), 15 reference(s)
 CODEN: LIFSAK ISSN: 0024-3205
 PUI S0024320500007840
 DT Journal; Article
 CY United States
 LA English
 SL English

L61 ANSWER 9 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN
 TI Purification of a novel apolipoprotein H-like milk protein with ribonucleolytic and cell-free translation inhibitory activities
 SO Life Sciences (2000), 67(8), 887-894
 CODEN: LIFSAK; ISSN: 0024-3205
 AU Ye, X. Y.; Ng, T. B.
 AN 2000:531051 HCAPLUS
 DN 133:234167

L61 ANSWER 10 OF 192 MEDLINE on STN DUPLICATE 4
 TI Ribonuclease, cell-free translation -inhibitory and superoxide radical scavenging activities of the iron-binding protein lactoferrin from bovine milk.
 SO The international journal of biochemistry & cell biology, (2000 Feb) Vol. 32, No. 2, pp. 235-41.
 Journal code: 9508482. ISSN: 1357-2725.
 AU Ye X Y; Wang H X; Liu F; Ng T B
 AN 2000150665 MEDLINE

L61 ANSWER 11 OF 192 MEDLINE on STN DUPLICATE 5
 TI Quinqueginsin, a novel protein with anti-human immunodeficiency virus, antifungal, ribonuclease and cell-free translation-inhibitory activities from American ginseng roots.
 SO Biochemical and biophysical research communications, (2000 Mar 5) Vol. 269, No. 1, pp. 203-8.
 Journal code: 0372516. ISSN: 0006-291X.
 AU Wang H X; Ng T B
 AN 2000160473 MEDLINE

L61 ANSWER 12 OF 192 MEDLINE on STN DUPLICATE 6
 TI Stabilization effect of zeolite on DHFR mRNA in a wheat germ cell -free protein synthesis system.
 SO Journal of bioscience and bioengineering, (2000) Vol. 89, No. 2, pp. 193-5.
 Journal code: 100888800. ISSN: 1389-1723.
 AU Jung G Y; Lee E Y; Kim Y; Jung B W; Kang S H; Choi C Y
 AN 2005557039 MEDLINE

L61 ANSWER 13 OF 192 MEDLINE on STN DUPLICATE 7
 TI Post-transcriptional regulation of rat CYP2E1 expression: role of CYP2E1 mRNA untranslated regions in control of translational efficiency and message stability.
 SO Archives of biochemistry and biophysics, (2000 Apr 1) Vol. 376, No. 1, pp. 180-90.
 Journal code: 0372430. ISSN: 0003-9861.
 AU Kocarek T A; Zangar R C; Novak R F
 AN 2000195423 MEDLINE

L61 ANSWER 14 OF 192 MEDLINE on STN DUPLICATE 8
 TI Dolichin, a new chitinase-like antifungal protein isolated from field beans (Dolichos lablab).
 SO Biochemical and biophysical research communications, (2000 Mar 5) Vol. 269, No. 1, pp. 155-9.
 Journal code: 0372516. ISSN: 0006-291X.
 AU Ye X Y; Wang H X; Ng T B
 AN 2000160466 MEDLINE

L61 ANSWER 15 OF 192 MEDLINE on STN DUPLICATE 9
 TI An easy cell-free protein synthesis system dependent on the addition of crude Escherichia coli tRNA.
 SO Journal of biochemistry, (2000 Jan) Vol. 127, No. 1, pp. 37-41.
 Journal code: 0376600. ISSN: 0021-924X.
 AU Kanda T; Takai K; Yokoyama S; Takaku H
 AN 2000198205 MEDLINE

L61 ANSWER 16 OF 192 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN
 TI Caspase-3 is the primary activator of apoptotic DNA fragmentation via DNA fragmentation factor-45/inhibitor of caspase-activated DNase inactivation.
 SO Journal of Biological Chemistry, (Oct. 22, 1999) Vol. 274, No. 43, pp. 30651-30656. print.
 CODEN: JBCHA3. ISSN: 0021-9258.
 AU Wolf, Beni B. [Reprint author]; Schuler, Martin; Echeverri, Fernando; Green, Douglas R.
 AN 2000:433616 BIOSIS

L61 ANSWER 17 OF 192 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN
 TI Ribonuclease activity of rat liver perchloric acid-soluble protein, a potent inhibitor of protein synthesis.
 SO Journal of Biological Chemistry, (July 16, 1999) Vol. 274, No. 29, pp.

20688-20692. print.

CODEN: JBCHA3. ISSN: 0021-9258.

AU Morishita, Ryo; Kawagoshi, Akihito; Sawasaki, Tatsuya; Madin, Kairat;
Ogasawara, Tomio; Oka, Tatsuzo; Endo, Yaeta [Reprint author]
AN 1999:467435 BIOSIS

L61 ANSWER 18 OF 192 MEDLINE on STN DUPLICATE 10
TI Functional differences of two forms of the inhibitor of
caspase-activated DNase, ICAD-L, and ICAD-S.
SO The Journal of biological chemistry, (1999 May 28) Vol. 274, No. 22, pp.
15740-4.
Journal code: 2985121R. ISSN: 0021-9258.
AU Sakahira H; Enari M; Nagata S
AN 1999269116 MEDLINE

L61 ANSWER 19 OF 192 MEDLINE on STN DUPLICATE 11
TI Inhibitors of DNA strand transfer reactions catalyzed by HIV-1 reverse
transcriptase.
SO Biochemistry, (1999 Oct 5) Vol. 38, No. 40, pp. 13070-6.
Journal code: 0370623. ISSN: 0006-2960.
AU Gabbara S; Davis W R; Hupe L; Hupe D; Peliska J A
AN 1999459252 MEDLINE

L61 ANSWER 20 OF 192 MEDLINE on STN DUPLICATE 12
TI Comparative inhibitory potential of differently modified antisense
oligodeoxynucleotides on hepatitis C virus translation.
SO European journal of clinical investigation, (1999 Oct) Vol. 29, No. 10,
pp. 868-76.
Journal code: 0245331. ISSN: 0014-2972.
AU Alt M; Eisenhardt S; Serwe M; Renz R; Engels J W; Caselmann W H
AN 2000051274 MEDLINE

L61 ANSWER 21 OF 192 LIFESCI COPYRIGHT 2006 CSA on STN DUPLICATE 13
TI Negative regulation of the pts operon by Mlc: mechanism underlying glucose
induction in Escherichia coli
SO Genes to Cells [Genes Cells], (19990700) vol. 4, no. 7, pp. 391-399.
ISSN: 1356-9597.
AU Tanaka, Y.; Kimata, K.; Inada, T.; Tagami, H.; Aiba, H.
AN 1999:111994 LIFESCI

L61 ANSWER 22 OF 192 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on
STN
TI Self-association of Puralpha is mediated by RNA.
SO Journal of Cellular Biochemistry, (Sept. 1, 1999) Vol. 74, No. 3, pp.
334-348. print.
CODEN: JCEBD5. ISSN: 0730-2312.
AU Gallia, Gary L.; Darbinian, Nune; Johnson, Edward M.; Khalili, Kamel
[Reprint author]
AN 1999:377823 BIOSIS

L61 ANSWER 23 OF 192 MEDLINE on STN DUPLICATE 14
TI In vivo and in vitro processing of the Bacillus subtilis transcript coding
for glutamyl-tRNA synthetase, serine acetyltransferase, and cysteinyl-tRNA
synthetase.
SO RNA (New York, N.Y.), (1999 Feb) Vol. 5, No. 2, pp. 281-9.
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PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
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WO 9423041	A3	19950105		
W:	AT, AU, BB, BG, BR, BY, CA, CH, CN, CZ, DE, DK, ES, FI, GB, HU, JP, KP, KR, KZ, LK, LU, LV, MG, MN, MW, NL, NO, NZ, PL, PT, RO, RU, SD, SE, SK, UA, UZ, VN			
RW:	AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE,			

	BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG	
CA 2159639	AA 19941013	CA 1994-2159639 19940401
AU 9464976	A1 19941024	AU 1994-64976 19940401
EP 693126	A1 19960124	EP 1994-912387 19940401
R: AT, BE, CH, DE, DK, ES, FR, GB, IT, LI, NL, SE		
JP 08509213	T2 19961001	JP 1994-522423 19940401

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 TI Protein preparation - by using cell-free protein synthesis system e.g. wheat germ extract.
 PI JP 06225783 A 19940816 (199437)* 5 C12P021-00
- L61 ANSWER 46 OF 192 SCISEARCH COPYRIGHT (c) 2006 The Thomson Corporation on STN
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 DN 122:96437

L61 ANSWER 54 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN
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 CODEN: PIXXD2
 IN Thompson, David V.; Van Oosbree, Thomas R.
 AN 1993:229729 HCAPLUS
 DN 118:229729

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 9307287	A1	19930415	WO 1992-US8518	19921007
W: AU, JP				
RW: AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LU, MC, NL, SE				
AU 9227921	A1	19930503	AU 1992-27921	19921007
AU 660329	B2	19950622		
EP 566714	A1	19931027	EP 1992-922343	19921007
EP 566714	B1	19970102		
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LI, LU, MC, NL, SE				
JP 06503477	T2	19940421	JP 1993-507150	19921007
JP 2904583	B2	19990614		
AT 147104	E	19970115	AT 1992-922343	19921007
ES 2097363	T3	19970401	ES 1992-922343	19921007

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 PI JP 05076381 A 19930330 (199317)* 4 C12P021-00

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 CODEN: PHPLAI; ISSN: 0031-9317
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 AN 1993:643690 HCAPLUS

DN 119:243690

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STN DUPLICATE 34
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phenol:chloroform extraction.
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ISSN: 0960-2585.
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Bewley, J. D.
AN 1993:432879 BIOSIS
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TI Characterization of ribonuclease H activities present in two cell-free
protein synthesizing systems, the wheat germ extract and the rabbit
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111-8.
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L61 ANSWER 65 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN
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translation system
SO PCT Int. Appl., 16 pp.
CODEN: PIXXD2
IN Ovodov, S. Yu.; Baranov, V. I.; Alakhov, Yu. B.; Ryabova, L. A.
AN 1991:554530 HCAPLUS
DN 115:154530

	PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
PI	WO 9102075	A1	19910221	WO 1990-SU145	19900605
	W: BG, CA, FI, HU, JP, US				
	RW: AT, BE, CH, DE, DK, ES, FR, GB, IT, LU, NL, SE				
	CA 2064685	AA	19910201	CA 1990-2064685	19900605
	CA 2064685	C	19960618		
	EP 485608	A1	19920520	EP 1990-912889	19900605
	EP 485608	B1	19951122		
	R: AT, BE, CH, DE, DK, ES, FR, GB, IT, LI, LU, NL, SE				
	JP 05505095	T2	19930805	JP 1990-512062	19900605
	JP 2891540	B2	19990517		
	AT 130633	E	19951215	AT 1990-912889	19900605
	US 5478730	A	19951226	US 1992-991757	19921216

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translation of matrix RNA modified by alkylation.
PI WO 9102074 A 19910221 (199110)*
RW: AT BE CH DE DK ES FR GB IT LU NL SE
W: BG CA FI HU JP US
IN ALAKHOV, J B; OVODOV, S J

L61 ANSWER 67 OF 192 MEDLINE on STN DUPLICATE 40
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1921-5.
Journal code: 2985121R. ISSN: 0021-9258.
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CODEN: NARHAD; ISSN: 0305-1048
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Nguyen Thanh Thuong; Toulme, Jean Jacques
AN 1991:507478 HCAPLUS
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NICOTIANA-ALATA (S-RNASE) ON INVITRO-GROWN POLLEN TUBES

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ISSN: 1040-4651.

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L61 ANSWER 71 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN

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CODEN: 59FGAZ

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DN 119:132433

L61 ANSWER 72 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 43

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SO PCT Int. Appl., 38 pp.

CODEN: PIXXD2

IN Balazs, Viktor; Balazs-Froehlich, Margit

AN 1991:554486 HCAPLUS

DN 115:154486

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 9009456	A1	19900823	WO 1990-DE102	19900216
W: JP, US				
RW: AT, BE, CH, DE, DK, ES, FR, GB, IT, LU, NL, SE				
EP 458831	A1	19911204	EP 1990-903145	19900216
R: AT, BE, CH, DE, DK, ES, FR, GB, IT, LI, LU, NL, SE				

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Journal code: 0370623. ISSN: 0006-2960.

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hepatoma cells is mediated by modification of RNA polymerase I or an
associated factor.

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1861-8.

Journal code: 8801431. ISSN: 0888-8809.

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L61 ANSWER 75 OF 192 LIFESCI COPYRIGHT 2006 CSA on STN

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Spodoptera frugiperda (Lepidoptera: Noctuidae) cells.

SO COMP. BIOCHEM. PHYSIOL., B., (1989) vol. 93B, no. 4, pp. 803-306.

AU Swerdel, M.R.; Fallon, A.M.

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DN 61:48966
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=> save temp l61 itt/a
ANSWER SET L61 HAS BEEN SAVED AS 'ITT/A'

=> s recbc? or exov or (exo or exonuclease) (w)v
FILE 'MEDLINE'
605 RECBC?
19 EXOV
4333 EXO
4163 EXONUCLEASE

442463 V
 93 (EXO OR EXONUCLEASE) (W) V
 L62 668 RECBC? OR EXOV OR (EXO OR EXONUCLEASE) (W) V

 FILE 'SCISEARCH'
 577 RECBC?
 11 EXOV
 10272 EXO
 3854 EXONUCLEASE
 1028711 V
 92 (EXO OR EXONUCLEASE) (W) V
 L63 612 RECBC? OR EXOV OR (EXO OR EXONUCLEASE) (W) V

 FILE 'LIFESCI'
 424 RECBC?
 12 EXOV
 2729 EXO
 2511 EXONUCLEASE
 79856 V
 46 (EXO OR EXONUCLEASE) (W) V
 L64 449 RECBC? OR EXOV OR (EXO OR EXONUCLEASE) (W) V

 FILE 'BIOTECHDS'
 59 RECBC?
 3 EXOV
 1314 EXO
 1222 EXONUCLEASE
 26520 V
 19 (EXO OR EXONUCLEASE) (W) V
 L65 76 RECBC? OR EXOV OR (EXO OR EXONUCLEASE) (W) V

 FILE 'BIOSIS'
 661 RECBC?
 21 EXOV
 8304 EXO
 4507 EXONUCLEASE
 336165 V
 100 (EXO OR EXONUCLEASE) (W) V
 L66 734 RECBC? OR EXOV OR (EXO OR EXONUCLEASE) (W) V

 FILE 'EMBASE'
 470 RECBC?
 17 EXOV
 4552 EXO
 3680 EXONUCLEASE
 416466 V
 76 (EXO OR EXONUCLEASE) (W) V
 L67 518 RECBC? OR EXOV OR (EXO OR EXONUCLEASE) (W) V

 FILE 'HCAPLUS'
 749 RECBC?
 26 EXOV
 28381 EXO
 6910 EXONUCLEASE
 1084227 V
 218 (EXO OR EXONUCLEASE) (W) V
 L68 901 RECBC? OR EXOV OR (EXO OR EXONUCLEASE) (W) V

 FILE 'NTIS'
 7 RECBC?
 0 EXOV
 296 EXO
 42 EXONUCLEASE
 39321 V
 2 (EXO OR EXONUCLEASE) (W) V

L69 9 RECBC? OR EXOV OR (EXO OR EXONUCLEASE) (W) V

FILE 'ESBIOBASE'

248 RECBC?

10 EXOV

1844 EXO

1776 EXONUCLEASE

261893 V

14 (EXO OR EXONUCLEASE) (W) V

L70 256 RECBC? OR EXOV OR (EXO OR EXONUCLEASE) (W) V

FILE 'BIOTECHNO'

354 RECBC?

17 EXOV

1171 EXO

2485 EXONUCLEASE

99304 V

39 (EXO OR EXONUCLEASE) (W) V

L71 373 RECBC? OR EXOV OR (EXO OR EXONUCLEASE) (W) V

FILE 'WPIDS'

14 RECBC?

0 EXOV

1910 EXO

928 EXONUCLEASE

274587 V

12 (EXO OR EXONUCLEASE) (W) V

L72 26 RECBC? OR EXOV OR (EXO OR EXONUCLEASE) (W) V

TOTAL FOR ALL FILES

L73 4622 RECBC? OR EXOV OR (EXO OR EXONUCLEASE) (W) V

=> s 124 and 173

FILE 'MEDLINE'

L74 4 L13 AND L62

FILE 'SCISEARCH'

L75 3 L14 AND L63

FILE 'LIFESCI'

L76 4 L15 AND L64

FILE 'BIOTECHDS'

L77 3 L16 AND L65

FILE 'BIOSIS'

L78 3 L17 AND L66

FILE 'EMBASE'

L79 3 L18 AND L67

FILE 'HCAPLUS'

L80 6 L19 AND L68

FILE 'NTIS'

L81 0 L20 AND L69

FILE 'ESBIOBASE'

L82 3 L21 AND L70

FILE 'BIOTECHNO'

L83 1 L22 AND L71

FILE 'WPIDS'

L84 1 L23 AND L72

TOTAL FOR ALL FILES

L85 31 L24 AND L73

=> dup rem l85

PROCESSING COMPLETED FOR L85

L86 7 DUP REM L85 (24 DUPLICATES REMOVED)

=> d tot

L86 ANSWER 1 OF 7 MEDLINE on STN DUPLICATE 1
 TI Increasing PCR fragment stability and protein yields in a cell-free system with genetically modified Escherichia coli extracts.
 SO Journal of molecular microbiology and biotechnology, (2005) Vol. 9, No. 1, pp. 26-34.
 Journal code: 100892561. ISSN: 1464-1801.
 AU Michel-Reydellet Nathalie; Woodrow Kim; Swartz James
 AN 2005576477 MEDLINE

L86 ANSWER 2 OF 7 MEDLINE on STN DUPLICATE 2
 TI A novel cell-free protein synthesis system.
 SO Journal of biotechnology, (2004 Jun 10) Vol. 110, No. 3, pp. 257-63.
 Journal code: 8411927. ISSN: 0168-1656.
 AU Sitaraman Kalavathy; Esposito Dominic; Klarmann George; Le Grice Stuart F; Hartley James L; Chatterjee Deb K
 AN 2004265272 MEDLINE

L86 ANSWER 3 OF 7 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN
 TI New in vitro protein or nucleic acid synthesis system comprising at least one extract from a cell, or inhibitor, and/or at least two energy sources providing chemical energy for synthesis, useful for producing proteins or nucleic acids;
 method for DNA synthesis or RNA synthesis
 AU CHATTERJEE D K; LONGO M C
 AN 2003-03912 BIOTECHDS
 PI WO 2002072890 19 Sep 2002

L86 ANSWER 4 OF 7 HCAPLUS COPYRIGHT 2006 ACS on STN
 TI Improvement of cell-free protein synthesis by modification of E. coli's genotype.
 SO Book of Abstracts, 219th ACS National Meeting, San Francisco, CA, March 26-30, 2000 (2000), BIOT-089 Publisher: American Chemical Society, Washington, D. C.
 CODEN: 69CLAC
 AU Michel-Reydellet, Nathalie; Kim, Dong-Myung; Swartz, James R.
 AN 2000:327311 HCAPLUS

L86 ANSWER 5 OF 7 MEDLINE on STN DUPLICATE 4
 TI Reconstitution of an SOS response pathway: derepression of transcription in response to DNA breaks.
 SO Cell, (1998 Dec 23) Vol. 95, No. 7, pp. 975-9.
 Journal code: 0413066. ISSN: 0092-8674.
 AU Anderson D G; Kowalczykowski S C
 AN 1999091055 MEDLINE

L86 ANSWER 6 OF 7 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN
 TI Novel plasmid and DNA;
 plasmid pIEPR12 vector containing a human cytomegalo virus immediate-early promoter regulatory region, for cloning in mammal cell culture or Escherichia coli
 AN 1993-01917 BIOTECHDS
 PI US 5168062 1 Dec 1992

L86 ANSWER 7 OF 7 MEDLINE on STN DUPLICATE 5

TI Enhanced polypeptide synthesis programmed by linear DNA fragments in
cell-free extracts lacking exonuclease V.
SO FEBS letters, (1983 Nov 14) Vol. 163, No. 2, pp. 221-4.
Journal code: 0155157. ISSN: 0014-5793.
AU Jackson M; Pratt J M; Holland I B
AN 84058336 MEDLINE

=> d ab 3-7

L86 ANSWER 3 OF 7 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN
AB DERWENT ABSTRACT:

NOVELTY - A new in vitro protein or nucleic acid synthesis system.

DETAILED DESCRIPTION - A new in vitro protein or nucleic acid synthesis system comprises one or more components selected from at least one extract from a cell having reduced activity of at least one enzyme that catalyzes hydrolysis of high energy phosphate bonds or hydrolysis or formation of phosphodiester bonds, at least one inhibitor of at least one enzyme that catalyzes hydrolysis of high energy phosphate bonds or hydrolysis or formation of phosphodiester bonds, and at least two energy sources providing chemical energy for synthesis. INDEPENDENT CLAIMS are also included for the following: (1) a composition comprising the one or more components cited above; (2) a kit for in vitro synthesis comprising the one or more components cited above; (3) a method for producing protein or nucleic acid from a nucleic acid template in an in vitro system; (4) a method for constructing an in vitro synthesis system; and (5) a composition comprising the one or more components cited above and at least one nucleic acid template in the presence of at least a partial synthesis product of the template.

BIOTECHNOLOGY - Preferred System: In the in vitro protein or nucleic acid synthesis system, at least one extract from a cell has reduced activity of at least one nuclease, phosphatase, or polymerase, and at least one inhibitor inhibits at least one nuclease, phosphatase or polymerase. The template is a DNA template. The nuclease is a DNase or RNase, preferably, exonuclease or endonuclease. The DNA endonuclease is endonuclease A. The RNA endonuclease is RNase E. The system further comprises at least one nucleic acid template, which is an RNA or DNA template. The system comprises at least one DNA template and an in vitro transcription/translation system. The phosphatase is an alkaline phosphatase. The energy sources generate or regenerate high-energy triphosphate compounds. The system comprises at least one or more compounds, as at least two different chemical fuel sources, selected from pyruvate, phosphoenolpyruvate (PEP), carbamoyl phosphate, acetyl phosphate, creatine phosphate, phosphopyruvate, glyceraldehydes-3-phosphate and glucose-6-phosphate. At least one extract from the cell is reduced in activity of at least one enzyme selected from OmpT, RNase E, alkaline phosphatase and endonuclease I. The system comprises further reduction in at least one activity of RNase I or RNase I asterisk. The enzyme can be RecBCD and the inhibitor can be at least Gam, preferably a soluble Gam. The system further comprises one or more nucleic acid templates or components such as an inhibitor of an enzyme that degrades the template, or at least one extract of a cell having reduced degradative effect on the template. The system comprises at least one energy source having at least two different energy sources, each of which generates or regenerates the high-energy triphosphate compounds for the synthesis. The two different chemical fuel sources comprise at least PEP and acetyl phosphate. The system can comprise at least one extract, nucleic acid template and energy source; at least one nucleic acid template and at least two energy sources; at least one extract; or at least one inhibitor. Preferred Kit: The kit further comprises one or more of the components selected from: (a) at least one inhibitor of RecBCD; (b) at least one cell, or its extract mutated at least one gene such as nuclease, polymerase or phosphatase; (c) an inhibitor of at least one enzyme such as nuclease, polymerase or phosphatase; (d) at least one energy source for synthesis; and (e) a

medium for growing of the at least one cell. Preferred Method: Producing protein or nucleic acid from the nucleic acid template in an in vitro system comprises: (a) contacting the template with at least one of the components cited above to form a mixture; and (b) incubating the mixture under conditions to produce at least one protein encoded by the template. Constructing an in vitro synthesis system comprises: (a) obtaining at least one cell extract; and (b) mixing the cell extract with the one or more components cited above. Preferred Composition: The product in the composition in (5) is a nucleic acid, preferably DNA or RNA.

USE - The system is useful for in vitro protein or nucleic acid synthesis (claimed). The extract from cell is useful for inhibiting or inactivating unwanted components/proteins/enzymes in the synthesis reaction. The inhibitors are useful for enhancing the production of the desired products in vitro. The kits are useful for facilitating the in vitro synthesis.

ADVANTAGE - The system provides a more efficient protein and nucleic acid synthesis by providing an improved energy supply for synthesis and by maintaining the nucleic acid templates for an extended synthesis reaction.

EXAMPLE - No relevant example given. (63 pages)

L86 ANSWER 4 OF 7 HCAPLUS COPYRIGHT 2006 ACS on STN

AB coli cell-free protein synthesis

systems have great potential but are limited by undesirable metabolic reactions. To improve these systems, we must first understand the limitations and then remove them. Knowledge of *E. coli* metabolism and genomic sequence as well as numerous genetic tools allows such anal. and optimization. Several targets have been identified. In phosphoenol pyruvate driven reactions, amino acid anal. showed that arginine, tryptophan and cysteine were being depleted. Preliminary expts. suggested that arginine decarboxylase (speA gene) and tryptophanase (tnaA gene) were responsible. One of our goals is to express proteins from PCR products. However, we observed that in cell exts. of the *E. coli* strain A19, linear DNA is unstable due to the activity of nucleases. Chief candidates are exonuclease V (recD gene) and endonuclease A (endA gene). These genes are targets for marker-free gene inactivation. The procedure for mutagenesis will be presented, as well as the impact on cell-free protein synthesis.

L86 ANSWER 5 OF 7 MEDLINE on STN

DUPLICATE 4

AB *E. coli* responds to DNA damage by derepressing the transcription of about 20 genes that make up the SOS pathway. Genetic analyses have shown that SOS induction in response to double-stranded DNA (dsDNA) breaks requires LexA repressor, and the RecA and RecBCD enzymes--proteins best known for their role as initiators of dsDNA break repair and homologous recombination. Here we demonstrate that purified RecA protein, RecBCD enzyme, single-stranded DNA-binding (SSB) protein, and LexA repressor respond to dsDNA breaks in vitro by derepressing transcription from an SOS promoter. Interestingly, derepression is more rapid if the DNA containing the dsDNA break has a chi recombination hot spot (5'-GCTGGTGG-3'), suggesting a novel regulatory role for one of the most overrepresented octamers in the *E. coli* genome.

L86 ANSWER 6 OF 7 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN

AB A new plasmid vector contains a specified DNA sequence from the immediate-early promoter regulatory region of human cytomegalo virus (HCMV), and a heterologous gene downstream, operatively linked. The sequence contains repeat sequences 19, 28, 16 and 21 bp in length. The plasmid is used to transform a eukaryote host, e.g. a mammal cell culture, or a microorganism, e.g. *Escherichia coli* RecBC-. The vector is preferably plasmid pIEPR12. The presence of HCMV repeat sequences facilitates transcription both in vitro and in vivo. In an example, in vitro transcription was tested with a HeLa cell lysate, and in vivo transcription was tested by transfection of human cells (by the calcium phosphate precipitate

method), extraction of cytoplasmic RNA and determination of the amount of specific RNA. Activation of an adjacent gene was at the transcriptional level, and transcription of the adjacent gene was increased by the presence of the viral sequences. (10pp)

L86 ANSWER 7 OF 7 MEDLINE on STN DUPLICATE 5
AB Preparation of an in vitro coupled transcription-translation system from E. coli strains lacking exonuclease V has greatly improved the system for use with added linear DNA fragments. In fact, in extracts of these mutants linear fragments are stable for several hours. However the cell extracts show a high level of endogenous background. To avoid this complication extracts were prepared at 30 degrees C from a mutant carrying a temperature-sensitive exonuclease V. Polypeptides coded by a specific DNA region, e.g., delineated by restriction endonuclease sites, can now be easily identified.

=> s linear dna

FILE 'MEDLINE'

174847 LINEAR

872961 DNA

L87 1414 LINEAR DNA
(LINEAR (W) DNA)

FILE 'SCISEARCH'

490252 LINEAR

620957 DNA

L88 1123 LINEAR DNA
(LINEAR (W) DNA)

FILE 'LIFESCI'

37376 "LINEAR"

283346 "DNA"

L89 939 LINEAR DNA
("LINEAR" (W) "DNA")

FILE 'BIOTECHDS'

7649 LINEAR

149142 DNA

L90 468 LINEAR DNA
(LINEAR (W) DNA)

FILE 'BIOSIS'

185492 LINEAR

1154202 DNA

L91 1611 LINEAR DNA
(LINEAR (W) DNA)

FILE 'EMBASE'

153306 "LINEAR"

661124 "DNA"

L92 1165 LINEAR DNA
("LINEAR" (W) "DNA")

FILE 'HCAPLUS'

585301 LINEAR

794147 DNA

L93 2114 LINEAR DNA
(LINEAR (W) DNA)

FILE 'NTIS'

73879 LINEAR

9431 DNA

L94 8 LINEAR DNA

(LINEAR (W) DNA)

FILE 'ESBIOBASE'
66041 LINEAR
295556 DNA
L95 595 LINEAR DNA
(LINEAR (W) DNA)

FILE 'BIOTECHNO'
25959 LINEAR
388151 DNA
L96 800 LINEAR DNA
(LINEAR (W) DNA)

FILE 'WPIDS'
237282 LINEAR
72504 DNA
L97 206 LINEAR DNA
(LINEAR (W) DNA)

TOTAL FOR ALL FILES
L98 10443 LINEAR DNA

=> s l24 and l98
FILE 'MEDLINE'
L99 40 L13 AND L87

FILE 'SCISEARCH'
L100 22 L14 AND L88

FILE 'LIFESCI'
L101 31 L15 AND L89

FILE 'BIOTECHDS'
L102 15 L16 AND L90

FILE 'BIOSIS'
L103 46 L17 AND L91

FILE 'EMBASE'
L104 32 L18 AND L92

FILE 'HCAPLUS'
L105 66 L19 AND L93

FILE 'NTIS'
L106 0 L20 AND L94

FILE 'ESBIOBASE'
L107 12 L21 AND L95

FILE 'BIOTECHNO'
L108 25 L22 AND L96

FILE 'WPIDS'
L109 8 L23 AND L97

TOTAL FOR ALL FILES
L110 297 L24 AND L98

=> s l110 not 2002-2006/py
FILE 'MEDLINE'
2844973 2002-2006/PY
(20020000-20069999/PY)
L111 34 L99 NOT 2002-2006/PY

FILE 'SCISEARCH'
5243521 2002-2006/PY
(20020000-20069999/PY)
L112 15 L100 NOT 2002-2006/PY

FILE 'LIFESCI'
500908 2002-2006/PY
L113 27 L101 NOT 2002-2006/PY

FILE 'BIOTECHDS'
122693 2002-2006/PY
L114 8 L102 NOT 2002-2006/PY

FILE 'BIOSIS'
2511100 2002-2006/PY
L115 37 L103 NOT 2002-2006/PY

FILE 'EMBASE'
2473613 2002-2006/PY
L116 28 L104 NOT 2002-2006/PY

FILE 'HCAPLUS'
5401054 2002-2006/PY
L117 52 L105 NOT 2002-2006/PY

FILE 'NTIS'
71750 2002-2006/PY
L118 0 L106 NOT 2002-2006/PY

FILE 'ESBIOBASE'
1474871 2002-2006/PY
L119 7 L107 NOT 2002-2006/PY

FILE 'BIOTECHNO'
244553 2002-2006/PY
L120 24 L108 NOT 2002-2006/PY

FILE 'WPIDS'
4666175 2002-2006/PY
L121 1 L109 NOT 2002-2006/PY

TOTAL FOR ALL FILES
L122 233 L110 NOT 2002-2006/PY

=> dup rem l122
PROCESSING COMPLETED FOR L122
L123 63 DUP REM L122 (170 DUPLICATES REMOVED)

=> d tot

L123 ANSWER 1 OF 63 HCAPLUS COPYRIGHT 2006 ACS on STN
TI Cell-free protein synthesis system
suitable for using PCR-amplified DNA templates: addition of excess
linear DNA
SO Jpn. Kokai Tokkyo Koho, 5 pp.
CODEN: JKXXAF
IN Sekiguchi, Akira; Arahata, Tomoya; Nunofuji, Satoshi; Nakano, Hideo;
Yamane, Tsuneo
AN 2001:365878 HCAPLUS
DN 134:363672
PATENT NO. KIND DATE APPLICATION NO. DATE

PI JP 2001136971 A2 20010522 JP 1999-323141 19991112

L123 ANSWER 2 OF 63 MEDLINE on STN DUPLICATE 1
 TI A novel mechanism controls anaerobic and catabolite regulation of the
Escherichia coli tdc operon.
 SO Molecular microbiology, (2001 Mar) Vol. 39, No. 5, pp. 1285-98.
 Journal code: 8712028. ISSN: 0950-382X.
 AU Sawers G
 AN 2001526297 MEDLINE

L123 ANSWER 3 OF 63 MEDLINE on STN DUPLICATE 2
 TI In-vitro competition analysis of procyclin gene and variant surface
 glycoprotein gene expression site transcription in *Trypanosoma brucei*.
 SO Molecular and biochemical parasitology, (2001 Mar) Vol. 113, No. 1, pp.
 55-65.
 Journal code: 8006324. ISSN: 0166-6851.
 AU Laufer G; Gunzl A
 AN 2001324191 MEDLINE

L123 ANSWER 4 OF 63 HCAPLUS COPYRIGHT 2006 ACS on STN
 TI Transcription in vitro using bacteriophage RNA
 polymerases
 SO Nucleic Acid Protocols Handbook (2000), 875-883. Editor(s): Rapley,
 Ralph. Publisher: Humana Press Inc., Totowa, N. J.
 CODEN: 68WSAO
 AU Schenborn, Elaine T.
 AN 2000:299241 HCAPLUS
 DN 134:66873

L123 ANSWER 5 OF 63 HCAPLUS COPYRIGHT 2006 ACS on STN
 TI Improvement of cell-free protein
 synthesis by modification of *E. coli*'s genotype.
 SO Book of Abstracts, 219th ACS National Meeting, San Francisco, CA, March
 26-30, 2000 (2000), BIOT-089 Publisher: American Chemical Society,
 Washington, D. C.
 CODEN: 69CLAC
 AU Michel-Reydellet, Nathalie; Kim, Dong-Myung; Swartz, James R.
 AN 2000:327311 HCAPLUS

L123 ANSWER 6 OF 63 MEDLINE on STN DUPLICATE 3
 TI Characterization of the *Moraxella catarrhalis* uspA1 and uspA2 genes and
 their encoded products.
 SO Journal of bacteriology, (1999 Jul) Vol. 181, No. 13, pp. 4026-34.
 Journal code: 2985120R. ISSN: 0021-9193.
 AU Cope L D; Lafontaine E R; Slaughter C A; Hasemann C A Jr; Aebi C;
 Henderson F W; McCracken G H Jr; Hansen E J
 AN 1999315796 MEDLINE

L123 ANSWER 7 OF 63 HCAPLUS COPYRIGHT 2006 ACS on STN
 TI Transcription in vitro using bacteriophage RNA
 polymerases
 SO Methods in Molecular Biology (Totowa, New Jersey) (1998), 86(RNA Isolation
 and Characterization Protocols), 209-219
 CODEN: MMBIED; ISSN: 1064-3745
 AU Schenborn, Elaine T.
 AN 1998:376273 HCAPLUS
 DN 129:145397

L123 ANSWER 8 OF 63 MEDLINE on STN DUPLICATE 4
 TI Elongation properties of vaccinia virus RNA polymerase: pausing, slippage,
 3' end addition, and termination site choice.
 SO Biochemistry, (1997 Dec 16) Vol. 36, No. 50, pp. 15892-9.
 Journal code: 0370623. ISSN: 0006-2960.
 AU Deng L; Shuman S
 AN 1998062306 MEDLINE

L123 ANSWER 9 OF 63 MEDLINE on STN DUPLICATE 5

TI DNA interactions of bifunctional dinuclear platinum(II) antitumor agents.
 SO European journal of biochemistry / FEBS, (1997 Jun 1) Vol. 246, No. 2, pp. 508-17.
 Journal code: 0107600. ISSN: 0014-2956.
 AU Zaludova R; Zakovska A; Kasparkova J; Balcarova Z; Kleinwachter V; Vrana O; Farrell N; Brabec V
 AN 97352550 MEDLINE

L123 ANSWER 10 OF 63 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN
 TI Direct expression of PCR products in a cell-free transcription/translation system: synthesis of antibacterial peptide cecropin; polymerase chain reaction direct in vitro transcription-translation system
 SO FEBS Lett.; (1997) 414, 2, 268-70
 CODEN: FEBLAL ISSN: 0014-5793
 AU Martemyanov K A; Spirin A S; *Gudkov A T
 AN 1997-11601 BIOTECHDS

L123 ANSWER 11 OF 63 LIFESCI COPYRIGHT 2006 CSA on STN DUPLICATE 6
 TI In vitro expression of PCR fragments: A convenient tool for generating purified recombinant proteins in analytical amounts
 SO Methods Mol. Cell. Biol., (19970000) vol. 6, no. 2, pp. 94-103.
 ISSN: 0898-7750.
 AU Nechansky, A.; Ruf, C.; Kricek, F.
 AN 1998:62355 LIFESCI

L123 ANSWER 12 OF 63 HCAPLUS COPYRIGHT 2006 ACS on STN
 TI Direct analysis of in vitro transcription products from super-coiled templates containing ribozymes
 SO BioTechniques (1996), 20(5), 750, 752, 754
 CODEN: BTNQDO; ISSN: 0736-6205
 AU Batt, David B.
 AN 1996:274055 HCAPLUS
 DN 124:334035

L123 ANSWER 13 OF 63 SCISEARCH COPYRIGHT (c) 2006 The Thomson Corporation on STN DUPLICATE 7
 TI In vitro expression of PCR fragments: A convenient tool for generating purified recombinant proteins in analytical amounts
 SO METHODS IN MOLECULAR AND CELLULAR BIOLOGY, (1995) Vol. 6, No. 2, pp. 94-103.
 ISSN: 0898-7750.
 AU Nechansky A (Reprint); Ruf C; Kricek F
 AN 1995:853495 SCISEARCH

L123 ANSWER 14 OF 63 MEDLINE on STN DUPLICATE 8
 TI Regulation of the Escherichia coli nrd operon: role of DNA supercoiling.
 SO Journal of bacteriology, (1994 Aug) Vol. 176, No. 15, pp. 4617-26.
 Journal code: 2985120R. ISSN: 0021-9193.
 AU Sun L; Fuchs J A
 AN 94321333 MEDLINE

L123 ANSWER 15 OF 63 MEDLINE on STN DUPLICATE 9
 TI A histone octamer can step around a transcribing polymerase without leaving the template.
 SO Cell, (1994 Jan 28) Vol. 76, No. 2, pp. 371-82.
 Journal code: 0413066. ISSN: 0092-8674.
 AU Studitsky V M; Clark D J; Felsenfeld G
 AN 94123343 MEDLINE

L123 ANSWER 16 OF 63 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN
 TI In vitro transcription of Clostridium acetobutylicum heat shock genes; possible role of inverted repeat in gene regulation of acetone and

butanol formation initiation (conference abstract)

SO Abstr.Gen.Meet.Am.Soc.Microbiol.; (1994) 94 Meet., 362
 CODEN: 0005P

AU Mueller H; Bahl H
 AN 1994-11192 BIOTECHDS

L123 ANSWER 17 OF 63 HCAPLUS COPYRIGHT 2006 ACS on STN
 TI Histone H1-mediated inactivation of methylated templates in vitro
 SO Biochemical Society Transactions (1994), 22(3), 298S
 CODEN: BCSTB5; ISSN: 0300-5127
 AU Johnson, Colin A.; Goddard, John P.; Adams, Roger L. P.
 AN 1994:572240 HCAPLUS
 DN 121:172240

L123 ANSWER 18 OF 63 MEDLINE on STN DUPLICATE 10
 TI DNA binding and bending are necessary but not sufficient for Fis-dependent
 activation of rrnB P1.
 SO Journal of bacteriology, (1993 Mar) Vol. 175, No. 6, pp. 1580-9.
 Journal code: 2985120R. ISSN: 0021-9193.
 AU Gosink K K; Ross W; Leirimo S; Osuna R; Finkel S E; Johnson R C; Gourse R L
 AN 93194781 MEDLINE

L123 ANSWER 19 OF 63 MEDLINE on STN DUPLICATE 11
 TI DNA topology and a minimal set of basal factors for transcription by RNA
 polymerase II.
 SO Cell; (1993 May 7) Vol. 73, No. 3, pp. 533-40.
 Journal code: 0413066. ISSN: 0092-8674.
 AU Parvin J D; Sharp P A
 AN 93258817 MEDLINE

L123 ANSWER 20 OF 63 WPIDS COPYRIGHT 2006 THE THOMSON CORP on STN
 TI Monoclonal antibodies to cruciform DNA structures - used for detection,
 isolation and study of cruciform DNA structures and associated gene
 elements.
 PI CA 1294906 C 19920128 (199308)* 26
 US 5192683 A 19930309 (199312)# 14 C12N005-20
 IN FRAPPIER, L; PRICE, G; ZANNIS-HADJOPOULOS, M

L123 ANSWER 21 OF 63 MEDLINE on STN DUPLICATE 12
 TI Differential effect of DNA supercoiling on transcription of
 adenovirus genes in vitro.
 SO The Journal of general virology, (1992 Oct) Vol. 73 (Pt 10), pp. 2631-8.
 Journal code: 0077340. ISSN: 0022-1317.
 AU Banerjee S; Spector D J
 AN 93019018 MEDLINE

L123 ANSWER 22 OF 63 MEDLINE on STN DUPLICATE 13
 TI The NusA and NusG proteins of Escherichia coli increase the in vitro
 readthrough frequency of a transcriptional attenuator preceding the gene
 for the beta subunit of RNA polymerase.
 SO The Journal of biological chemistry, (1992 Jan 25) Vol. 267, No. 3, pp.
 1449-54.
 Journal code: 2985121R. ISSN: 0021-9258.
 AU Linn T; Greenblatt J
 AN 92112854 MEDLINE

L123 ANSWER 23 OF 63 MEDLINE on STN DUPLICATE 14
 TI Application of the polymerase chain reaction to the ribonuclease
 protection assay.
 SO BioTechniques, (1992 Dec) Vol. 13, No. 6, pp. 922-7.
 Journal code: 8306785. ISSN: 0736-6205.
 AU Yang H; Melera P W
 AN 93119653 MEDLINE

L123 ANSWER 24 OF 63 MEDLINE on STN DUPLICATE 15

TI DNA supercoiling response of the sigma 54-dependent *Klebsiella pneumoniae* nifL promoter in vitro.
 SO Journal of molecular biology, (1992 Jun 5) Vol. 225, No. 3, pp. 591-607.
 Journal code: 2985088R. ISSN: 0022-2836.
 AU Whitehall S; Austin S; Dixon R
 AN 92292148 MEDLINE

L123 ANSWER 25 OF 63 MEDLINE on STN DUPLICATE 16
 TI Defining nucleic acid-binding properties of avian retrovirus integrase by deletion analysis.
 SO Journal of virology, (1991 Mar) Vol. 65, No. 3, pp. 1160-7.
 Journal code: 0113724. ISSN: 0022-538X.
 AU Mumm S R; Grandgenett D P
 AN 91140704 MEDLINE

L123 ANSWER 26 OF 63 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN
 TI A coupled in vitro transcription-translation system for the exclusive synthesis of polypeptides expressed from the T7 promoter;
 high level gene expression in *Escherichia coli* cell-free extract
 SO FEBS Lett.; (1991) 291, 2, 259-63
 CODEN: FEBLAL
 AU Nevin D E; *Pratt J M
 AN 1991-14515 BIOTECHDS

L123 ANSWER 27 OF 63 SCISEARCH COPYRIGHT (c) 2006 The Thomson Corporation on STN
 TI PATCHED CIRCLE POLYMERASE CHAIN-REACTION
 SO BIOTECH FORUM EUROPE, (MAR 1991) Vol. 8, No. 3, pp. 130-132.
 ISSN: 0938-7501.
 AU SONG C Z (Reprint); YANG K Y
 AN 1991:426701 SCISEARCH

L123 ANSWER 28 OF 63 MEDLINE on STN DUPLICATE 18
 TI Effects of the DNA topoisomerase II inhibitor, VM26, on transcriptional initiation in vitro.
 SO Life sciences, (1990) Vol. 46, No. 18, pp. 1309-18.
 Journal code: 0375521. ISSN: 0024-3205.
 AU Preston G M; White B A
 AN 90265212 MEDLINE

L123 ANSWER 29 OF 63 MEDLINE on STN DUPLICATE 19
 TI Transcription elongation factor SII (TFIIS) enables RNA polymerase II to elongate through a block to transcription in a human gene in vitro.
 SO The Journal of biological chemistry, (1989 Jun 25) Vol. 264, No. 18, pp. 10799-809.
 Journal code: 2985121R. ISSN: 0021-9258.
 AU Reines D; Chamberlin M J; Kane C M
 AN 89278155 MEDLINE

L123 ANSWER 30 OF 63 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN
 TI A novel yeast secretion signal isolated from 28K killer precursor protein encoded on the linear DNA plasmid pGKL1;
 foreign protein secretion and glycosylation from *Saccharomyces cerevisiae* host
 SO Nucleic Acids Res.; (1988) 16, 15, 7499-511
 CODEN: NARHAD
 AU Tokunaga M; Wada N; Hishinuma F
 AN 1988-10612 BIOTECHDS

L123 ANSWER 31 OF 63 LIFESCI COPYRIGHT 2006 CSA on STN DUPLICATE 20
 TI Difference between supercoiled and linear deoxyribonucleic acids in preinitiation complex formation for accurate transcription in vitro.

SO CHEM. PHARM. BULL. (TOKYO)., (1988) vol. 36, no. 7, pp. 2517-2522.
AU Ohtsuki, M.; Nakanishi, Y.; Sekimizu, K.; Natori, S.
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L123 ANSWER 32 OF 63 MEDLINE on STN DUPLICATE 21
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L123 ANSWER 1 OF 63 HCAPLUS COPYRIGHT 2006 ACS on STN
AB Described is an improved cell-free protein synthesis system using PCR amplified DNA as template. Addition of excess linear nucleic acids unrelated to the template in the reaction mixture to counteract the degradation of linear DNA template is described. Cell-free synthesis of chloramphenicol acetyl transferase (CAT) using Escherichia coli cell extract; T-7 RNA polymerase; and a DNA construct containing T7 promoter, the ribosome-binding site, the template DNA, and the T7 terminator, with addition of salmon sperm DNA or calf thymus DNA is described.

L123 ANSWER 4 OF 63 HCAPLUS COPYRIGHT 2006 ACS on STN
AB The authors provide detailed methods for synthesis of RNA by transcription in vitro from a linear DNA template. Preparation of the DNA template, the transcription reaction, and enrichment of the RNA product are all discussed.

L123 ANSWER 5 OF 63 HCAPLUS COPYRIGHT 2006 ACS on STN
AB coli cell-free protein synthesis systems have great potential but are limited by undesirable metabolic reactions. To improve these systems, we must first understand the limitations and then remove them. Knowledge of E. coli metabolism and genomic sequence as well as numerous genetic tools allows such anal. and optimization. Several targets have been identified. In phosphoenol pyruvate driven reactions, amino acid anal. showed that arginine, tryptophan and cysteine were being depleted. Preliminary expts. suggested that arginine decarboxylase (speA gene) and tryptophanase (tnaA gene) were responsible. One of our goals is to express proteins from PCR products. However, we observed that in cell exts. of the E. coli strain A19, linear DNA is unstable due to the activity of nucleases. Chief candidates are exonuclease V (recD gene) and endonuclease A (endA gene). These genes are targets for marker-free gene inactivation. The procedure for mutagenesis will be presented, as well as the impact on cell-free protein synthesis.

L123 ANSWER 7 OF 63 HCAPLUS COPYRIGHT 2006 ACS on STN
AB The authors describe the synthesis of RNA by transcription from a linear DNA template using bacteriophage RNA polymerases.

L123 ANSWER 10 OF 63 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN
AB A simple method was developed for direct expression of polymerase chain

reaction (PCR)-generated linear DNA in a cell-free transcription-translation system, without cloning fragments in a plasmid vector. The method was adapted to production of recombinant cecropin from an artificial gene. An oligonucleotide encoding cecropin was synthesized, and a set of tandem multimeric genes without stop codons was obtained by modified splicing overlap extension. A megaprimer containing a phage T7 promoter and ribosome binding site, with s10 leader and Shine-Dalgarno sequences, was amplified by PCR from plasmid pET21d(+), and used to add expression elements to the gene. An unpurified PCR mixture was added to *Escherichia coli* S30 extract with phage T7 RNA-polymerase (EC-2.7.7.6) for cell-free transcription-translation, giving a cecropin yield of up to 3 nmol monomer/ml. This approach should be useful for direct in vitro expression of polypeptides and proteins that are unstable in living cells, or proteins that are strongly cytotoxic. (14 ref)

L123 ANSWER 11 OF 63 LIFESCI COPYRIGHT 2006 CSA on STN DUPLICATE 6

AB A method is presented in which an in vitro gene expression system for linear DNA templates based on bacterial extracts was directly coupled to a protein purification method. This method allows the production of purified, biologically active, recombinant proteins directly from polymerase-chain-reaction-amplified DNA containing all regulatory elements necessary for transcription and translation in a one-tube reaction within a few hours. The purified material can be directly used in biological testing. Because in vitro transcription or translation is carried out in bacterial extracts, subsequent large-scale production of the desired recombinant protein can be performed in *Escherichia coli* by using the same construct as for the in vitro expression, thus avoiding laborious and time-consuming subcloning. Furthermore, by using this system, host cell transformation and growth, in vivo gene expression, and protein purification procedures can be avoided for many applications.

L123 ANSWER 12 OF 63 HCAPLUS COPYRIGHT 2006 ACS on STN

AB Currently there are two methods to analyze promoter activities in vitro. The first method involves transcription from supercoiled plasmids. To reduce the time of this assay, one can use a linearized DNA template instead of the supercoiled template. This second method involves incubating the linearized DNA, mammalian extract, NTPs and a radiolabeled ribonucleotide triphosphate. This produces a radiolabeled runoff transcript. This method has the disadvantage of being less efficient because linear DNA is transcribed less efficiently than supercoiled DNA. This dilemma was overcome by the (G)-free transcription cassette. Although this technique works very well, it is not under natural conditions because GTP is absent. Ribozymes have allowed a novel approach to this problem. Here the authors describe a method that allows the use of a supercoiled template and all four NTPs, utilizing a self-cleaving hammerhead ribozyme.

L123 ANSWER 13 OF 63 SCISEARCH COPYRIGHT (c) 2006 The Thomson Corporation on STN DUPLICATE 7

AB A method is presented in which an in vitro gene expression system for linear DNA templates based on bacterial extracts was directly coupled to a protein purification method. This method allows the production of purified, biologically active, recombinant proteins directly from polymerase-chain-reaction-amplified DNA containing all regulatory elements necessary for transcription and translation in a one-tube reaction within a few hours. The purified material can be directly used in biological testing. Because in vitro transcription or translation is carried out in bacterial extracts, subsequent large-scale production of the desired recombinant protein can be performed in *Escherichia coli* by using the same construct as for the in vitro expression, thus avoiding laborious and time-consuming subcloning. Furthermore, by using this system, host cell transformation

and growth, in vivo gene expression, and protein purification procedures can be avoided for many applications. Methods Mol. Cell. Biol. 6:94-103, 1997. (C) 1997 Wiley-Liss, Inc.

- L123 ANSWER 21 OF 63 MEDLINE on STN DUPLICATE 12
AB We examined the effect of DNA template topology on the transcription of immediate early (E1a), early (E1b) and late (pIX) adenovirus genes in vitro. Transcription in whole cell extracts was measured by quantitative hybridization to end-labelled DNA and protection of hybrids from S1 nuclease digestion. Two- to fourfold more E1a RNA was synthesized from supercoiled, compared to linear, DNA templates. Similarly, transcription of the E1b gene was stimulated three- to sevenfold when the template was supercoiled. In contrast, RNA synthesis from the late pIX gene was found to be independent of DNA topology. These results show that DNA topology affects transcription in a promoter-specific manner.
- L123 ANSWER 26 OF 63 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN
AB A coupled transcription-translation in vitro system was developed in Escherichia coli for the generation of polypeptides cloned downstream of the phage T7 promoter. The system consists of a cell-free, DNA-free extract of E. coli S30 (cells contain endogenous T7 RNA-polymerase, EC-2.7.7.6), rifampicin (an E. coli RNA-polymerase-inhibitor), a labeled amino acid (usually 35S-methionine) and a low mol.weight mixture which provides a supplement of factors including amino acids and nucleotide triphosphates. When primed with template DNA (phage, plasmid or linear DNA fragments), preferably plasmid pMJ216, encoding the target gene under exclusive control of the T7 promoter, this system has the capability to synthesize relatively large amounts of a unique, labeled polypeptide. This coupled system is simple to prepare and use, and yields more polypeptide product from a given amount of DNA template than when the processes of transcription and translation are uncoupled. (13 ref)
- L123 ANSWER 31 OF 63 LIFESCI COPYRIGHT 2006 CSA on STN DUPLICATE 20
AB A nuclear extract of Ehrlich ascites tumor cells was separated into three fractions that were essential for accurate transcription initiation. Studies with these fractions showed that the efficiency of transcription of supercoiled deoxyribonucleic acid (DNA) was better than that of linear DNA. A clear difference was found in the rates of formation of a preinitiation complex with these two templates. Consistent with this difference, the results of gel shift assay showed that the complex of supercoiled DNA with transcription factors is much more complicated than that of linear DNA.
- L123 ANSWER 32 OF 63 MEDLINE on STN DUPLICATE 21
AB In a posterior silk gland extract, covalently closed circular (ccc) DNA is in a superhelical state that supports more transcription of fibroin gene than does linear DNA. A HeLa cell extract showed neither the supercoiling activity nor the preference for the transcription of ccc DNA over linear DNA. These activities could be added to the HeLa cell extract. Phosphocellulose fractionation of the posterior silk gland extract yielded a flow-through fraction and a 0.6 M KCl eluate fraction that were required for the supercoiling and for the efficient transcription of the ccc template in the acceptor HeLa cell extract. The 0.6 M KCl fraction had a DNA topoisomerase II activity, and the flow-through fraction contained a supercoiling factor that, with the aid of topoisomerase II, introduced negative supercoils into ccc DNA. When both fractions were added to the posterior silk gland extract, more supercoiling occurred than with the extract alone. Several genes were optimally transcribed under various extents of supercoiling. The fibroin gene and adenovirus 2 major late promoter were fully transcribed as partially supercoiled templates. The sericin gene required more supercoiling for full transcription, whereas no preference for supercoiling was seen with the transcription of hsp70. These results

suggest that DNA topology plays a role in the regulation of gene expression.

L123 ANSWER 33 OF 63 HCAPLUS COPYRIGHT 2006 ACS on STN

AB The constraints imposed on DNA by torsional stress and the consequences of resulting DNA topol. on in vitro transcription were studied. Form V DNA, prepared by annealing complementary circular DNA single strands of pBR322, with a resulting superhelical d. greater than 10 times that observed in vivo, was used to demonstrate that approx. half of the DNA sites were resistant to methylation and altered in structure. Very short stretches of alternate and normal structure were observed. Measurements of DNA rotation on the surface of nucleosomes and nucleosome rotation with respect to each other revealed that adjacent nucleosomes in chromatin cannot rotate relative to each other and that isolated topol. domains exist at each nucleosome. Systematic variation of an in vitro system for transcription of SV40 DNA resulted in conditions where at low DNA concns. (2.5 µg/mL) transcription from supercoiled DNA was maximal while that from linear DNA was insignificant, demonstrating a clear in vitro effect of DNA topol. on the level of gene expression. Comparison of both linear and supercoiled plasmid DNA by pulsed gel electrophoresis reveals that supercoils reorient about 10 times faster in a gel than the corresponding linear forms, implying that the average supercoil condenses the net mol. length by 10-fold. The aspects of such condensed and torsionally stressed DNA on eukaryotic gene expression is discussed.

L123 ANSWER 37 OF 63 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN

AB The use of DNA-dependent protein synthesizing systems for studying the mechanisms of translation, transcription and their regulation is reviewed. The need for a simpler in vitro coupled system is a result of the development of recombinant DNA technology and the increasing use of plasmids. Accordingly, partially reconstituted systems were described for studies on ribosomal protein genes and the problem of linear DNA expression was discussed. Highly defined systems containing up to 30 different, purified protein factors required for transcription and translation are a means of obtaining new information in a system free from interference of gene expression by contaminants. A method which would enable the production of single proteins in a transcription-translation system of prokaryotic and eukaryotic origin is regarded as a new approach towards the development of a universal in vitro expression system. (105 ref)

L123 ANSWER 44 OF 63 MEDLINE on STN

DUPLICATE 28

AB Transcription of a tRNAArg-tRNAAsp gene pair from *Saccharomyces cerevisiae* by an homologous yeast extract results in a dimeric precursor molecule which is processed to mature-sized tRNAArg and tRNAAsp molecules. We have transcribed linear DNA fragments cleaved within the gene sequences to show that precursor synthesis is not dependent on the internal promoter of the second gene (tRNAAsp). Furthermore, the second gene does not support independent transcription when the normal upstream initiation site is removed.

L123 ANSWER 45 OF 63 HCAPLUS COPYRIGHT 2006 ACS on STN

AB The transcription activity of an in vitro mouse RNA polymerase system with double-stranded calf thymus DNA as template was increased 2-4- and 1-3-fold by the addition of 1-10 mM spermine (I) and spermidine (II), resp. At concns. of 0.2-0.5 mM, I also stimulated the transcription activity of an in vitro *Escherichia coli* RNA polymerase system in the presence of double-stranded DNA; however I at concns. of 1.0-5.0 mM markedly reduced the transcription activity. II activated transcription activity in the *E. coli* RNA polymerase system at all concns. tested (0.2-5.0 mM). The effect of polyamines, especially I, on transcription activity of both RNA polymerase systems was dependent on the template DNAs used. In the presence of a double-stranded DNA template, RNA synthesis in the mouse RNA polymerase system was stimulated by I at

all concns. (0.5-10 mM) tested, whereas RNA synthesis was not affected by I when single-stranded denatured calf thymus DNA or single-stranded circular DNA (M13 phage) was used as template. With the double-stranded DNA template, ≤ 0.5 mM I stimulated RNA synthesis in the E. coli RNA polymerase system, but inhibited RNA synthesis at ≥ 1 mM. The E. coli RNA synthesis system was inhibited by all concns. of I tested in the presence of single-stranded linear DNA template, but was slightly enhanced when the single-stranded circular DNA template was used.

- L123 ANSWER 46 OF 63 MEDLINE on STN DUPLICATE 29
AB A recB21 derivative (CLB7) of an Escherichia coli rna-19 pnp-7 strain (PR7) was constructed for use in examining the in vitro coupled transcription-translation of linear DNA. The expression of linearized DNAs in CLB7 (recB21 rna-19 pnp-7) lysates was enhanced significantly when compared with expression of the same DNAs in lysates prepared from the PR7 or the original recB21 (CF300) strains. In addition, the endogenous incorporation of [35S]methionine into protein was considerably reduced in CLB7 lysates relative to lysates derived from the original recB21 strain.
- L123 ANSWER 50 OF 63 MEDLINE on STN DUPLICATE 33
AB Preparation of an in vitro coupled transcription-translation system from E. coli strains lacking exonuclease V has greatly improved the system for use with added linear DNA fragments. In fact, in extracts of these mutants linear fragments are stable for several hours. However the cell extracts show a high level of endogenous background. To avoid this complication extracts were prepared at 30 degrees C from a mutant carrying a temperature-sensitive exonuclease V. Polypeptides coded by a specific DNA region, e.g., delineated by restriction endonuclease sites, can now be easily identified.
- L123 ANSWER 52 OF 63 MEDLINE on STN DUPLICATE 35
- L123 ANSWER 53 OF 63 MEDLINE on STN DUPLICATE 36
AB Heretofore the DNA-directed coupled transcription-translation system, most useful in gene expression analysis, has been limited to the use of circular or long linear DNAs. Linear DNAs are degraded in this system by an exonucleolytic activity that can be eliminated by making the synthetic extracts from a suitable recB mutant of Escherichia coli. Using these extracts, we have examined the gene expression of a variety of linear DNAs. In particular, the complex pattern of expression of ribosomal protein genes and RNA polymerase genes in the rpoBC-rplLJ region has been analyzed by comparing the protein products obtained when using lambda rifd18 DNA with the product obtained when using the same DNA segmented with various restriction enzymes. The results obtained confirm the conclusions of others obtained by much more elaborate in vivo techniques. It seems highly likely that this cell-free system will have extensive applications in the area of analysis of gene expression.
- L123 ANSWER 58 OF 63 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN DUPLICATE 40
AB Several steps in the transcription of double-stranded polyoma DNA by calf thymus RNA polymerase II(B) were analyzed. EM observations show that, as found for the Escherichia coli enzyme, the eucaryotic RNA polymerase exhibits a higher affinity for superhelical DNA as compared to linear DNA. The half-time of complexes formed at 37° C between the enzyme and superhelical DNA is relatively short (30-60 s), compared with 30 h for the E. coli enzyme under identical conditions. Binary complexes formed with linear DNA have a faster dissociation rate. The same results were obtained by using the nitrocellulose binding assay or by measuring the decay of heparin-resistant complexes. Only ternary complexes, enzyme-DNA-RNA, obtained after the formation of the 1st phosphodiester bond, are resistant

to polyanions. As for the procaryotic enzyme, RNA polymerase II probably has to melt locally the double-stranded DNA during initiation, but it probably lacks a σ type factor necessary to stabilize the open initiation complexes. RNA polymerase II synthesizes shorter RNA chains than E. coli RNA polymerase. The formation of RNase I resistant RNA-DNA hybrids during transcription may explain the reduced rate of chain elongation.

L123 ANSWER 59 OF 63 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN DUPLICATE 41

AB EM of nascent RNA chains was used to localize promoters on linear and negatively superhelical [phage] λ DNA. Transcription was performed in vitro using Escherichia coli RNA polymerase. RNA from 4 promoters was seen on linear λ DNA: these include the λ PL and PR promoters, which are the main early λ promoters in vivo, and 2 promoters within the b2 region. To orient the circular DNA for EM, a restriction enzyme isolated from Streptomyces albus G (Sal I) was used to cleave the DNA at unique points. The Sal I cleavage sites on λ DNA were at 67.3% and 68.3% on the linear map. Visualization of nascent RNA transcribed from superhelical λ DNA indicates that transcription increases from PL and PR with a particularly large increase from PL, and there is a transcription activation from promoters that are not used on the linear DNA and that coincide with the areas of λ which are most readily denatured and which possess the highest A + T content. The activation of such regions and the increased efficiency of the promoters used on linear DNA are discussed in terms of the energetics of unwinding a negatively superhelical DNA by a ligand such as RNA polymerase. The association of A + T-rich regions with promoters and the possible role of superhelicity in transcriptional activation in vivo are discussed.

L123 ANSWER 62 OF 63 HCAPLUS COPYRIGHT 2006 ACS on STN

AB Polypeptides made in an in vitro DNA-dependent protein-synthesizing system from the linear DNAs of different mutants and hybrids of bacteriophage λ were examined. The methionine-35S-labeled products were analyzed by Na dodecyl sulfate gel electrophoresis and by 2-dimensional fingerprinting of their tryptic peptides. Anal. of promoter mutants (sex1 and x13), of amber mutants in gene N, of deletion mutants (λ b2, λ db30-7 nin5), and of substitution derivs. λ i434, λ i21 hy5) permitted assignment of many peptides to definite portions of λ DNA. Similar anal. of purified DNA fragments, produced by digestion of λ DNA with endonuclease RI (R.N. Yoshimori, 1971), made it possible to exam. independently the products controlled by different promoters. Most of the coupled polypeptide synthesis arose from early regions of λ DNA in vitro. Effects of repressor and of the transcription termination factor rho were also tested. Both efficiently reduced the over-all level of coupled translation, and with a specificity that helped characterize some gene products. The use of bacterial components from λ -infected or uninfected cells resulted in the same pattern of polypeptides. No difference was detected when circular instead of linear DNA mols. were used as template.

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L61 ANSWER 12 OF 192 MEDLINE on STN DUPLICATE 6

AB The effects of zeolites and monocations on the protein synthesis in a cell-free system derived from wheat germ were investigated. M type of synthetic zeolite markedly enhanced the translation efficiency. Whereas this kind of stimulatory effect of zeolite in an Escherichia coli cell-free system resulted from a change in the salt compositions of the reaction solution with the addition of zeolite, the enhancement of protein synthesis in a wheat germ cell-free system was not due to the ion exchange reaction of zeolites. From the results of mRNA stability analysis, it was found that zeolite could stabilize the mRNA in a wheat germ cell-free protein synthesis system. The stabilization of mRNA by the simple addition of zeolites is useful for the enhancement of protein synthesis in a wheat germ cell-free system, since conventional methods to improve mRNA stability, such as the addition of nuclease inhibitor, have not been effective for a wheat germ cell-free system.

L61 ANSWER 24 OF 192 SCISEARCH COPYRIGHT (c) 2006 The Thomson Corporation on STN DUPLICATE 15

AB Degradation of template mRNA lowers the efficiency of cell-free translation. Although cell-free translation system using wheat germ extract (WGE) is usually supplemented with placental RNase inhibitor (PRI) to stabilize mRNA, the decay rate of mRNA, however, was still fairly rapid. For elucidating cause of mRNA degradation in wheat germ cell-free translation system, the authors applied activity staining in SDS-PAGE gel to investigate ribonuclease activities in WGE. RNases or nucleases of molecular mass of about 65, 42, and 37 kDa were detected in WGE, and named WGa, WGb and WGC, respectively. WGb and WGC were nucleases that degraded both RNA and DNA, and WGa was an RNase. Polyguanylic acid (5'), poly[G], which was known to have affinity with RNases, strongly inhibited RNase activity in WGE. Addition of poly[G] to wheat germ cell-free translation system resulted in 7-fold increase in protein productivity. Luciferase mRNA that was added in the cell-free translation system as a template was not degraded in 90 min in the presence of poly[G] while it was degraded by 80% in 30 min in the absence of poly[G]. The authors concluded that poly[G] improved the protein productivity of wheat germ cell-free translation system by inhibiting RNase activity that degrades template RNA. (C) 1999 Elsevier Science B.V. All rights reserved.

L61 ANSWER 29 OF 192 MEDLINE on STN DUPLICATE 20

AB mRNA stability is a limiting parameter for the efficiency of in vitro protein biosynthesis. In order to develop strategies to prolong the mRNA half-life, we investigated the ribonuclease activities in the complete Escherichia coli system, in the separate cell fractions 70S ribosomes and S-100 and in the non-cellular fraction. Our results imply that the amount

of ribonucleolytic activities and the insensitivity to placental RNase inhibitor in the complete system are due to the 70S ribosome fraction, whereas the generation of small degradation products is due to the S-100 fraction. Remarkably, the human placental RNase inhibitor is able to reduce mRNA degradation in the bacterial S-100 fraction.

L61 ANSWER 31 OF 192 SCISEARCH COPYRIGHT (c) 2006 The Thomson Corporation
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AB Antisense oligonucleotides, modified by a manganese(III)tris(4-N-methylpyridiniumyl)porphyrin (an artificial nuclease) and activated by an oxidant to promote their nuclease activity, exhibit a higher inhibitory effect on the in vitro translation of chloramphenicol acetyltransferase (CAT) mRNA than the corresponding unmodified oligonucleotides. This effect is as high as that observed in the presence of E. Coli RNase H.

L61 ANSWER 35 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN

AB Cell-free protein synthesis using wheat germ extract in the conventional batch reaction was studied, focusing on both a higher protein synthesis rate and a longer synthesis period. The following points were improved with considerable success: (1) Optimization of the reaction conditions of cell-free protein synthesis enabled a longer protein synthesis, ≤ 10 h, even in a batch system. (2) Condensation of wheat germ extract by precipitation with polyethylene glycol increased the initial protein synthesis rate by approx. 10-fold; protein yield reached 0.12 mg/mL in only 3 h with the use of Cu^{2+} , a possible inhibitor of RNase and phosphatase.

L61 ANSWER 36 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN

AB Copper ion, an inhibitor to RNase, 0.1-1 mM is used for in vitro protein synthesis using cell extract. Protein synthesis using Escherichia coli extract in the presence of copper ion was shown.

L61 ANSWER 43 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN

AB In the Escherichia coli cell-free system, modification of the cell extract can be achieved by preparation of strains carrying an addnl. property or of those being induced with a certain gene expression prior to harvesting. The authors analyzed the cell-free system with S30 extract containing T7 RNA polymerases (S30 extract-T7pol) prepared from an E. coli BL21(DE3) strain, which includes T7 RNA polymerase from extrinsic genes by IPTG induction, as a model for the improvement of the cell-free system. The fact that a significant degree of mRNA degradation was observed in the cell-free system with S30 extract-T7pol indicates that an increase in RNase activity was an unfavorable influence derived from the cell-extract modification process. This influence was prohibited by the addition of an effective RNase inhibitor, such as copper ion, to the reaction mixture

L61 ANSWER 45 OF 192 WPIDS COPYRIGHT 2006 THE THOMSON CORP on STN

AB JP 06225783 A UPAB: 19941109

Preparation of protein by using a cell-free protein synthesis system containing cell-free extract in which the protein concentrate is 15 mg/ml. or lower, the cell-free extract being pref. wheat germ extract.

USE/ADVANTAGE - Protein is prepared efficiently ca. 3 to 6 times higher than the conventional method.

In an example, wheat germ extract was prepared according to the method of K. Maruc and B. Dudock. 5 ml. of the crude extract was ultrafiltered through M10 to give 1 ml of wheat germ extract. The reaction liquor consisted of 20 mM HEPES buffer, 1 mM ATP, 20 micro MGTP. 2 mM DTF, 4mM

creatin phosphate, 40 micro-g/ml. creatine phosphokinase, 0.1 mM spermidine, 0/01 mM spermine, 40 micro-M amino acid, 0.5 mCi/ml. (35S) methionine, 1.0 U/micro-l RNase inhibitor, 75 ng/micro-l BMV RNA, 70 mM K acetate, 2 mM Mg acetate and 2 micro-l wheat germ extract and the total volume was made to be 10 micro-l. The reaction was carried out at 30deg.C. for 1 hr.. The protein concentration originated

from

the cell-free extract in the reaction liquor was 15 mg/ml. compared to 5 mg/ml. for a control using the crude extract.

Dwg.0/0

L61 ANSWER 54 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN

AB A method for coupling transcription and translation from DNA using a modified eukaryotic cell-free extract containing Mg salts at a concentration that

allows both transcription and translation occur in the system. The concentration

of Mg is 2.5.apprx.3.5 mM, preferably, 2.6.apprx.3.0 mM. Transcription and translation of the luciferase gene from a DNA construct using a solution containing standard rabbit reticulocyte lysate, rNTP's, SP6 RNA polymerase, RNasin, Mg acetate, and other ingredients were demonstrated.

L61 ANSWER 65 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN

AB A cell-free translation system for manufacture of proteins contains ribosomes, matrix RNA, ATP, GTP, and amino acids. The products are removed as they are formed, and ATP, GTP, and amino acids are continually added to maintain their original concentration. The system is immobilized in microcapsules with membranes of polyelectrolyte complexes. Thus, 32,000 pmol of calcitonin was synthesized over 100 h using a wheat lysate, ribosomes, mRNA, and substrates immobilized in Na alginate-poly-L-lysine microcapsules.

L61 ANSWER 75 OF 192 LIFESCI COPYRIGHT 2006 CSA on STN

AB Conditions for in vitro translation of mRNA in cell-free extracts from cultured *Spodoptera frugiperda* cells were defined. Incorporation of (super(35)S)methionine into acid-precipitable material increased for approximately 1 hr, and was sensitive to the protein synthesis inhibitors pactamycin and cycloheximide. Micrococcal nuclease-treated lysate, primed with purified rabbit globin mRNA, synthesized a major protein with the size of full length globin, indicating that the lysate supported correct initiation and elongation of polypeptides.

L61 ANSWER 88 OF 192 MEDLINE on STN DUPLICATE 54

AB A phosphocellulose flowthrough fraction required for accurate transcription in vitro by RNA polymerase II was found to contain a DNase inhibitor which was necessary to maintain template integrity (Price D.H., Sluder A.E. & Greenleaf A.L. (1987) J. Biol. Chemical 262, 3244-3255). Starting with a *Drosophila* Kc cell nuclear extract, the DNase inhibitory activity has been purified 19,000-fold. In combination with the other necessary fractions, the highly purified inhibitor continues to support reconstruction of transcription. It thus appears to be the only required activity in the original phosphocellulose flowthrough fraction. The inhibitor is a protein which does not bind to DNA or inhibit DNase I, so that it has also been useful in assays for DNA binding proteins in crude, DNase-contaminated fractions.

L61 ANSWER 91 OF 192 MEDLINE on STN DUPLICATE 56

AB Cell-free protein synthesis in rabbit reticulocyte lysate translation mixtures has been studied during multi-hour incubations. In an impaired lysate obtained from cells stored at 0 degrees C before lysis, and showing a low initial rate of synthesis, translation could be stimulated by a factor of 4 by including RNase inhibitor and additional ATP and GTP. In

translation mixtures prepared from normal lysates, protein synthesis could be improved by approximately 50% by the addition of excess GTP. The observed increases in protein synthesis were obtained by improved maintenance of the initial rate of synthesis.

L61 ANSWER 96 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN

AB A method is described for the preparation of wheat germ exts. for use in an in vitro translation system for the synthesis of high-mol.-weight proteins by modifying the method of W. Zagorski (1978) and by using high spermidine concns., human placental RNase inhibitor, and mRNAs extracted from plant RNA viruses. The efficiency of the method is comparable to that obtained in the reticulocyte lysate system.

L61 ANSWER 108 OF 192 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN DUPLICATE 66

AB Three inhibitors of animal cell-free protein synthesis were isolated from barley seeds by ammonium sulfate fractionation. CM-cellulose chromatography and gel filtration on Bio-Gel P-60. The 3 inhibitors were all single chain, basic proteins with MW .apprx. 31,000 and with very similar amino acid compositions. In the rabbit reticulocyte lysate system, they inhibited in vitro translation to 50% at concentrations between 15 and 25 ng . ml-1, while in the wheat germ system, their effect was very weak. The 3 translation inhibitors showed no detectable RNase activity neither on ribosomal RNA from rabbit liver nor on homopolynucleotides.

L61 ANSWER 115 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN

AB Modified procedures for preparation of rough microsomes (RM) and membrane-bound polysomes from rat liver that are active in cell-free protein synthesis in vitro are described. The major modifications are: (1) cell homogenization is performed in a slightly hypertonic sucrose solution lacking salts to prevent aggregation of organelles; (2) postmitochondrial supernatant is used as a source of the RM fraction, potent RNase inhibitor, and endogenous salts to keep microsomes in contact with the supernatant throughout the preparation; and (3) pelleting of the RM fraction is avoided by using sucrose cushions.

L61 ANSWER 140 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN

AB The S-30 fraction (30,000 g supernatant) prepared from Escherichia coli by the method of Zubay et al. was treated with various concns. of DNase I at 30° for 30 min. At 0.1 µg/mL, DNase I decreased the rate of endogenous DNA-dependent protein formation by 90%. The DNase inhibitor G-actin at 50 µg/mL antagonized the effect of DNase I (0.5 µg/mL). G-actin alone had no effect on protein formation. This system may be used in the study of in vitro protein synthesis.

L61 ANSWER 144 OF 192 MEDLINE on STN DUPLICATE 84

L61 ANSWER 146 OF 192 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN

L61 ANSWER 153 OF 192 MEDLINE on STN DUPLICATE 91

AB The RNA species synthesized in vitro by a transcribing nucleoprotein (TNP) complex of vesicular stomatitis virus (VSV) were translated with high efficiency in a fractionated cell-free system derived from reticulocytes. The use of TNP complexes isolated from VSV Indiana, VSV New Jersey, and Chandipura viruses showed that in each case the predominant polypeptides synthesized had electrophoretic mobilities identical to their virion N, NS, and M polypeptides in proportions reflecting those found in infected cells rather than purified virions. A minor polypeptide corresponding to unglycosylated polypeptide G was also observed, but the in vitro synthesis

of polypeptide L was not detected. The addition of RNase inhibitor to transcription mixtures markedly increased the rate of RNA synthesis. Furthermore, the messenger activity of the RNA was significantly enhanced. The inclusion of S-adenosyl L-methionine during transcription substantially increased the messenger activity of the product RNA, suggesting a requirement for methylation. Fractionation by oligodeoxythymidylic acid-cellulose chromatography revealed that the RNA required a polyadnylic acid tract for messenger activity.

L61 ANSWER 169 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN

AB DNA hydrolysis in chromatin by DNase I (EC 3.1.4.5) or II (EC 3.1.4.6) remained limited even after prolonged incubation. This limitation might be due to a redistribution under the action of DNase of f1 histone on chromatin DNA. With large amts. of DNase, all of the chromatin DNA may, however, be digested. A comparison was made between the extent of DNA hydrolysis by DNase and the resulting inhibition of chromatin template activity for endogenous, Escherichia coli, or rat liver (form II) RNA polymerases. The repressed sequences did not seem to be less sensitive to DNase action than the derepressed, since the decrease in template activity parallels rather well the digestion of DNA.

L61 ANSWER 185 OF 192 HCAPLUS COPYRIGHT 2006 ACS on STN

AB cf. preceding abstract DNA from T-phages (T2, T4 am82, and T3), Bacillus subtilis phage SP50, B. subtilis, B. megaterium, and Haemophilus influenzae actively initiated DNA-dependent protein synthesis in E. coli cell-free systems. DNA from E. coli and the replicative form of phage ϕ X 174 were moderately active. DNA from E. coli phages λ c and λ vir, from polyoma and papilloma viruses, and from calf thymus, calf liver Ehrlich ascites tumor cells, and regenerating rat liver were practically inactive. All of the DNA preps. were able to initiate the DNA dependent RNA synthesis in the presence of E. coli RNA polymerase. The hydrolysis of messenger RNA synthesized in the cell-free system by endogenous ribonucleases was inhibited by yeast ribosomal RNA. The undesirable inhibition of the transcription process could be avoided if the transcription was begun before the addition of the RNA. The stability of RNA and protein synthesis in the cell-free system was considerably increased by the addition of a ribonuclease inhibitor (probably a protein of nucleoprotein) from rabbit reticulocyte supernatant. If the DNA from E. coli phage T4 was used as the template, amino acids were incorporated into proteins which precipitated with antibodies against early protein.

L61 ANSWER 189 OF 192 MEDLINE on STN

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DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)	SINCE FILE	TOTAL
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